

NOTICE OF DRAFTSPERSON'S PATENT DRAWING REVIEW

1-20-99

1. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Lichtenthaler and Whistler (1973). The total chlorophyll content was determined by the method of Arar and Cook (1980). The carotenoid content was determined by the method of Lichtenthaler and Whistler (1973). The total carotenoid content was determined by the method of Arar and Cook (1980). The total protein content was determined by the method of Lowry et al. (1951). The total lipid content was determined by the method of Bligh and Dyer (1959). The total carbohydrate content was determined by the method of Dubois and Gilles (1950). The total nucleic acid content was determined by the method of Burton (1956). The total ash content was determined by the method of AOAC (1990). The total moisture content was determined by the method of AOAC (1990). The total dry matter content was determined by the method of AOAC (1990). The total organic acid content was determined by the method of AOAC (1990). The total alkaloid content was determined by the method of AOAC (1990). The total saponin content was determined by the method of AOAC (1990). The total tannin content was determined by the method of AOAC (1990). The total flavonoid content was determined by the method of AOAC (1990). The total phenol content was determined by the method of AOAC (1990). The total terpenoid content was determined by the method of AOAC (1990). The total steroid content was determined by the method of AOAC (1990). The total glycoside content was determined by the method of AOAC (1990). The total alkaloid content was determined by the method of AOAC (1990). The total saponin content was determined by the method of AOAC (1990). The total tannin content was determined by the method of AOAC (1990). The total flavonoid content was determined by the method of AOAC (1990). The total phenol content was determined by the method of AOAC (1990). The total terpenoid content was determined by the method of AOAC (1990). The total steroid content was determined by the method of AOAC (1990). The total glycoside content was determined by the method of AOAC (1990).

By using the `mean` function, we can calculate the means indicated by μ . The parameter `na.rm = TRUE` tells the function to ignore missing values. The `sd` function calculates the standard deviations indicated by σ . The parameter `na.rm = TRUE` tells the function to ignore missing values. The `var` function calculates the variances indicated by σ^2 . The parameter `na.rm = TRUE` tells the function to ignore missing values. The `cov` function calculates the covariances indicated by σ_{12} and σ_{21} . The parameter `na.rm = TRUE` tells the function to ignore missing values. The `cor` function calculates the correlations indicated by ρ_{12} and ρ_{21} . The parameter `na.rm = TRUE` tells the function to ignore missing values. The `plot` function is used to create a scatter plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `summary` function is used to get a summary of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `str` function is used to get the structure of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `table` function is used to get the frequency table of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `hist` function is used to get the histogram of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `boxplot` function is used to get the boxplot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `qqnorm` function is used to get the Q-Q plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `qqline` function is used to get the Q-Q line of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `acf` function is used to get the ACF plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `pacf` function is used to get the PACF plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `arma` function is used to get the ARMA model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `ar` function is used to get the AR model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `ma` function is used to get the MA model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `arima` function is used to get the ARIMA model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `forecast` function is used to get the forecast of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `plot` function is used to create a plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `summary` function is used to get a summary of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `str` function is used to get the structure of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `table` function is used to get the frequency table of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `hist` function is used to get the histogram of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `boxplot` function is used to get the boxplot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `qqnorm` function is used to get the Q-Q plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `qqline` function is used to get the Q-Q line of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `acf` function is used to get the ACF plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `pacf` function is used to get the PACF plot of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `arma` function is used to get the ARMA model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `ar` function is used to get the AR model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `ma` function is used to get the MA model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `arima` function is used to get the ARIMA model of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values. The `forecast` function is used to get the forecast of the data. The parameter `na.rm = TRUE` tells the function to ignore missing values.

[illegible]

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